SEQUENCE LISTING

æ	(1) GENERAL INFORMATION:
5	(i) APPLICANT:
	(A) NAME: University of Guelph(B) STREET: Gordon St.(C) CITY: Guelph
10	(D) STATE: Ontario (E) COUNTRY: Canada
	(F) POSTAL CODE (ZIP): NIG 2W1
15	(ii) TITLE OF INVENTION: Monocot Transformation Using Agrobacterium
	(iii) NUMBER OF SEQUENCES: 4
20	(iv) COMPUTER READABLE FORM:(A) MEDIUM TYPE: Floppy disk(B) COMPUTER: IBM PC compatible
	(C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
25	(v) CURRENT APPLICATION DATA:
	APPLICATION NUMBER: FILING DATE:
	(2) INFORMATION FOR SEQ ID NO: 1:
30	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs
	(B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear
35	(ii) MOLECULE TYPE: other nucleic acid
	(A) DESCRIPTION: /desc = "primer 1"
40	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
45	CCGTCTGCGG GAGCGCTATC C 21
	(2) INFORMATION FOR SEQ ID NO: 2:
50	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 21 base pairs(B) TYPE: nucleic acid
	(C) STRANDEDNESS: single (D) TOPOLOGY: linear
55	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "primer 2"</pre>
JJ	(A) Diochar Levi. Acoc - primer 2

5	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:	
J	CATCGCAAGA CCGGCAACAG G	21
	(2) INFORMATION FOR SEQ ID NO: 3:	
10	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 18 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
15	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc ≈ "primer 3"</pre>	
20		
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:	
25	AGAAACCAAA GGGTCCTG	18
20	(2) INFORMATION FOR SEQ ID NO: 4:	
30	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
35	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "primer 4"</pre>	
40	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:	
	03.003.03.000. 3.00003.00	10